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Lotus japonicus mRNA for cytochrome P450, complete cds. AB024931
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophy
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots;
Rosidae, eurosids I, Fabales, Fabaceae, Papilionoideae, Loteae;
                                                                                                            1641 GTTCCAAGGGCACATAACCTCATCTGTGTCCGGTTGCAAGATCAAGTGCCGTACCCAAA 1700
                                   Lotus japonicus (strain:Gifu) root cDNA to mRNA, clone:LjCYP-1 Lotus japonicus
                                                                                                                                155 GTTCCAAGGGCACATAACCTCATCTGTGTCCCGGTTGCAAGATCAAGTGCCGTACCCAAA 214
                                                       Induction of isoflavonoid pathway in the model legume Lotus japonicus: molecular characterization of enzymes involved in phytoalexin biosynthesis
35 ATGGCCACACTGCTTGCGTCAGTTATCCAGTGCTTTGATCTCAGCGTAGTGGGCCCCACAG 94
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/note="Lotus corniculatus var. japonicus"
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Shimada, N., Akashi, T., Aoki, T. and Ayabe, S.
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Shimada,N., Aoki,T. and Ayabe,S.
Direct Submission
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Cloning of cytochrome P450 cDNAs from cultured Glycyrrhiza echinata
L. cells and their transcriptional activation by elicitor-treatment
Plant Sci. 126, 39-47 (1997)
Location/Qualifiers
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Glycyrrhiza echinata
Glycyrrhiza echinata
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Galegeae;
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Akashi,T., Aoki,T., Takahashi,T., Kameya,N., Nakamura,I. and
                             GATCAAGTGCCGTACCCAAACTCTTTTCGTCGTAAAACATACGCGCGACACCAGAAAGCT
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Organisma-"Glycyrrhiza echinata"
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Direct Submission
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TREQUESALRENTYDNSVAMVPPAPTVMKETRIKTYNNLINPLLNATTVNKLEPLRSQEIRKVL
KRAMAGSABSOKPLNATEBLLKWTNSTISRMALGBAEHYKDIVREVLKIFGERSLIPRI
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ATAGMATLLSSVIQCFELQVVGPKGQILKGSDAKANMEERRAGLTVPRANSLMCVPLAR ö Gaps ô Length 1557; Indels 1.8%; Score 35; DB 8; I 100.0%; Pred. No. 2.7e-08; 390 GCCICCACCCTGAACTITTCAAACTCTTCCTCCA 424 0; Mismatches 385 365 g SKVAAELLSS 366 c 35; Conservative ò

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ATAGMATLLSSVIQCFELOVVGPKGQILKGSDAKANMEERAGLTVPRANSLMCVPLAR
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VGHIALDOPLANSLIKIGERYGPLYTPEGSMPTVVASTPETEKRETOTHEATSFS
TREOGSAIRRLTYDNSVAAVPPEAPTWETIRKYIMNDLINATTVNKLRPLASQEIRKVL
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EVDSVVGKDRLVDESDIQNLPYIRAIVKETFRMHPPLPVVKRKCVQECELNGYVIPEG
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Shimada, N., Aoki, T. and Ayabe, S.
Direct Submission
Direct Submission
Dispartment of Applied Biological Science; Rameino 1866, Fujisawa,
Ranagawa 252-8510, Japan (E-mail:ayabe@brs.nihon-u.ac.jp,
Tel:81-466-84-3703, Fax:81-466-80-1141)
Location/Qualifiers
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Lotus japonicus
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophy
Spermatophyta; Magnollophyta; eudicotyledons; core eudicots;
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                                 GTTCCAAGGGCACATAACCTCATCTGTGTCCGGTTGCAAGATCAAGTGCCGTACCCAAA 1700
                                                 35 ATGGCCACACTCCTTGCGTCAGTTATCCAGTGCTTTGATCTCAGCGTAGTGGGCCCACAG 94
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Shimada,N., Akashi,T., Aoki,T. and Ayabe,S.
Induction of isoflavonoid pathway in the model legume Lotus
japonicus: molecular characterization of enzymes involved in
phytoalexin blosynthesis
Plant science (Shannon, Ireland) 160 (1), 37-47 (2000)
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100.0%; Pred. No. 2.7e-08;
ive 0; Mismatches 0; Indels
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Location/Qualifiers
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
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Direct Submission
Direct Submission
Submitted (20-NOV-1996) Shin-ichi Ayabe, Nihon University,
Department of Applied Biological Science; Kameino 1866, Fujisawa,
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                       1621 AAGAGAGGTGGACTCACGGTTCCAAGGCACATAACCTCATCTGTGTCCCGGTTGCAA 1680
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Glycyrrhiza echinata suspension-cultured cells cDNA to mRNA,
clone_lib:lambda ZapII clone:Ge-8.
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/clone_lib="lambda ZapII"
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